



PATENT COOPERATION TREATY

To:

From the INTERNATIONAL BUREAU

PCT

NOTIFICATION OF ELECTION

(PCT Rule 61.2)

United States Patent and Trademark Office (Box PCT)

Crystal Plaza 2 Washington, DC 20231 ETATS-UNIS D'AMERIQUE

Date of mailing (day/month/year) 04 August 1997 (04.08.97)	in its capacity as elected Office
International application No. PCT/NO96/00266	Applicant's or agent's file reference Eij 1 HV
International filing date (day/month/year) 13 November 1996 (13.11.96)	Priority date (day/month/year) 13 November 1995 (13.11.95)
Applicant	
EIJSINK, Vincent, G., H. et al	

1	The designated Office is hereby notified of its election made:	
	X in the demand filed with the International Preliminary Examining Authority on:	
	03 June 1997 (03.06.97)	
	in a notice effecting later election filed with the International Bureau on:	
2	The election X was	
	made before the expiration of 19 months from the priority date or, where Rule 32 applies, within the time limit under Rule 32.2(b).	

The International Bureau of WIPO 34, chemin des Colombettes 1211 Geneva 20, Switzerland Authorized officer

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	tates of America
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International application No. PCT/NO 96/00266

A. CLASSIFICATION OF SUBJECT MATTER

IPC6: C12N 15/74, C07K 14/335
According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC6: C12N, C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

SE,DK,FI,NO classes as above

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

WPI, FULLTEXT, MEDLINE, BIOSIS, DERWENT BIOTECH ABS, EMBL/GENBANK/ SWISSPROT/DDBJ

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	Molecular Microbiology, Volume 18, No 4, 1995, Dzung Bao Diep et al, "A bacteriocin-like peptide induces bacteriocin synthesis in Lactobacillus plantarum C11" page 631 - page 639	1-15,17-21
х	Microbiology, Volume 140, 1994, Petra S. Tichaczek et al, "Cloning and sequencing of sakP encoding sakacin P, the bacteriocin produced by Lactobacillus sake LTH 673", page 361 - page 367, see especially fig 2	1-15,17-21
		

	Further documents are listed in the continuation of Box	C.	X See patent family annex.		
*	Special categories of cited documents:		later document published after the international filing date or priority		
"A"	document defining the general state of the art which is not considered to be of particular relevance		date and not in conflict with the application but cited to understand the principle or theory underlying the invention		
″E″	erlier document but published on or after the international filing date		document of particular relevance: the claimed invention cannot be		
"L"	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other		considered novel or cannot be considered to involve an inventive step when the document is taken alone		
	special reason (as specified)		document of particular relevance: the claimed invention cannot be		
-0-	document referring to an oral disclosure, use, exhibition or other means		considered to involve an inventive step when the document is combined with one or more other such documents, such combination		
~P~	document published prior to the international filing date but later than		being obvious to a person skilled in the art		
l	the priority date claimed	"& "	document member of the same patent family		
Dat	e of the actual completion of the international search	Date of	mailing of the international search report		
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13	February 1997				
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Swe	edish Patent Office				
Box 5055, S-102 42 STOCKHOLM			Patrick Andersson		
Fac	simile No. +46 8 666 02 86	Telephone No. +46 8 782 25 00			



International application No. PCT/NO 96/00266

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C (Continu	nation). DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the rele	evant passages	Relevant to claim No.
х	Molecular Microbilogy, Volume 17, No 3, 1995, Koen Venema et al, "Functional analysis o pediocin operon of Pediococcus acidilacti PAC1.0: PedB is the immunity protein and the precursor processing enzyme", page 515 - page 522, fig 1 and summary	f the ci	1-15,17-21
x	Applied and EnvironmentalMicrobiology, Volume No 1, January 1994, Dzung Bao Diep et al Gene Encoding Plantaricin A, a Bacterioci Lactobacillus plantarum C11, Is Located o Same Transcription Unit as an agr-Like Re System" page 160 - page 166	, "The n from n the	1-15,17-21
х	Journal of Bacteriology, Volume 177, No 8, Ap 1995, Lars Axelsson et al, "The Genes Inv Production of and Immunity to Sakacin A, Bacteriocin from Lactobacillus sake Lb706 page 2125 - page 2137	olved in a	1-2,6-15, 17-21
x	Applied and Environmental Microbiology, Volum No 2, February 1991, Marco J. van Belkum "Organization and Nucleotide Sequences of Lactococcal Bacteriocin Operons", page 492 - page 498, figure 1	et al,	1-2,6-15, 17-21
x	EP 0493779 A1 (QUEST INTERNATIONAL B.V.), 8 July 1992 (08.07.92), page 3, line 24 -	line 31	1-2,6-15, 17-21
х	WO 9404682 A1 (DZIEGLEWSKA, HANNA, EVA), 3 March 1994 (03.03.94), page 11, line 12 - line 37		1-2,6-15, 17-21
х	WO 119802 A1 (HOLMES, MICHAEL, J.), 25 December 1991 (25.12.91), page 6, line 9 - line 36		1-2,6-15, 17-21
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D.

Inter...uional application No.

PCT/NO 96/00266

Patent document cited in search report		Publication date		Patent family member(s)		
EP-A1-	0493779	08/07/92	AU-B- AU-A- CA-A- JP-A- US-A- US-A-	638616 8820091 2056086 7067652 5175252 5260212	01/07/93 09/07/92 01/07/92 14/03/95 29/12/92 09/11/93	
WO-A1-	9404682	03/03/94	AU-A-	4968893	15/03/94	
WO-A1-	119802	25/12/91	NONE			

PATENT COOPERATION TREAT PEC'D

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INTERNATIONAL PRELIMINARY EXAMINATION REPORT

			(PCT Article 36	and Hule	9 70)	·
Applicant's or	agent's	file reference	FOR FURTHER AC	TION	See	Notification of Transmittal of International minary Examination Report (PCT/IPEA/416)
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International a	pplicati	on No.	International filing date (day/i	month/year)		Priority date (day/month/year)
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1. This into	ernatio	onal preliminary exa	mination report has been pro t according to Article 36.	epared by ti	113 1111	emationary rollinary
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IV		Lack of unity of inv	/ention			the standard and annicability.
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VI		Certain document				
VII			the international application			
VIII		Certain observation	ons on the international appl	ication		
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INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/NO96/00266

I. Basi	s of the	report
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1. This report has been drawn on the basis of (substitute sheets which have been furnished to the receiving Office in response to an invitation under Article 14 are referred to in this report as "originally filed" and are not annexed to the report since they do not contain amendments.):

	resp the r	eport since they o	do not contain amendme	ents.):
	Des	cription, pages:		
	1-27		as originally filed	
	Clai	ms, No.:		
	1-15		with telefax of	16/02/1998
	Drav	wings, sheets:		
	1/3-	3/3	as originally filed	
2.	The	amendments hav	ve resulted in the cance	llation of:
		the description,	pages:	
		the claims,	Nos.:	
		the drawings,	sheets:	
3.		This report has to considered to go	been established as if (so beyond the disclosure	some of) the amendments had not been made, since they have been as filed (Rule 70.2(c)):
4.	Add	ditional observatio	ons, if necessary:	

INTERNATIONAL PRELIMINARY EXAMINATION REPORT

International application No. PCT/NO96/00266

- V. Reasoned statement under Article 35(2) with regard to novelty, inventive step or industrial applicability; citations and explanations supporting such statement
- 1. Statement

Novelty (N)

Yes:

Claims 1-15

No:

Claims

Inventive step (IS)

Yes:

Claims 1-15

No:

Claims

Industrial applicability (IA)

Yes:

Claims 1-15

No:

Claims

2. Citations and explanations

see separate sheet

SECTION	٧	

- The present application relates to an inducible gene expression system in lactic acid bacteria. It has been discovered that expression of the IF-K-R gene cluster is autoinduced by the gene product of the IF gene.
- 2. The quoted documents are:
 - D1: P. Tichaczek et al., Microbiology, Vol. 140, pp. 361-367, 1994
 - D2: Axelsson et al., Journal of Bacteriology, Vol. 177, No. 8, pp. 2125-2137, 1995
- 3. D1 discloses the sequence of bacteriocin SakP, its promoter and two further open reading frames (orfX and orfY), whose function is unknown. It is speculated that these ORFs could encode immunity proteins.
 - D2 is concerned with an operon containing the genes necessary for production and immunity of the bacteriocin sakacin A (SapA) from *Lactobacillus sake* Lb706. It is assumed that proteins SapK and SapR constitute a two-component signal-transduction system, mediating a response to an environmental signal, while SapT and SapE seem to be involved in secretion of SapA. SaiA encodes the immunity peptide. Two further open reading frames were identified (Orf4 and Orf1) but no function has been accorded to them. Both are similar to the bacteriocin leader peptides and are speculated to be structural genes for bacteriocins.
- 4. Claim 1 relates to a gene expression system in lactic acid bacteria which is characterised in that it comprises a gene/genes of interest which have been operably linked by genetic engineering to a strongly regulated promoter whose activity can be induced by an unmodified peptide. The promoter elements and the

INTERNATIONAL PRELIMINARY International application No. PCT/NO96/00266 EXAMINATION REPORT - SEPARATE SHEET

peptide are defined in the claim by all essential elements thereof. It is also specified in the claim that the genes of interest are **not identical** to the genes that are operably linked to said promoter elements in the lactic acid bacteria from which said promoter elements are derived.

The latter feature distinguishes the claimed gene expression system from naturally occurring lactic acid bacteria which inherently comprise the promoter elements specified in the claim.

A gene expression system as defined in Claim 1 has neither been disclosed nor rendered obvious in any of the cited documents. The subject-matter of Claim 1 and of the claims which are directly or indirectly dependent thereon is therefore considered novel and also appears to involve the required inventive step (Article 33(2) and (3) EPC).

5. Claim 11 relates to a peptide which is defined by its amino acid sequence. The peptide is capable of activating the inducible promoter described in Claim 1. Said peptide has not been disclosed in the prior art and its sequence could not have been derived plainly and logically from any of the cited documents. Novelty and inventive step are therefore recognized for the subject-matter of Claim 11 (Article 33(2) and (3) PCT).

CLAIMS

- 1. Gene expression system, characterized in that it comprises genes, promoter sequences and peptides involved in the production of bacteriocins except nisin in lactic acid bacteria.
- 2. Gene expression system, characterized in that it contains at least one specific regulated promoter, genes involved in transducing signals that induce gene expression, a peptide being that signal, and, possibly, genes involved in producing, processing, and secreting this inducing peptide.
- 3. The expression system of claims 1 and 2, characterized in that the said peptide is capable of inducing its own production and/or that of one or more bacteriocins by the lactid acid bacteria.
- 4. The expression system of claim 3, characterized in that said peptide is a functional analogue of the peptide of claim 3, functional analogues being defined as shortened, enlarged or mutated variants that retain the potential to induce gene expression.
- 5. The expression system of claim 4, wherein said peptide has the sequence Met-Ala-Gly-Asn-Ser-Ser-Asn-Phe-Ile-His-Lys-Ile-Lys-Gln-Ile-Phe-Thr-His-Arg, (seq. id. no. 1) or has the sequence Lys-Ser-Ser-Ala-Tyr-Ser-Leu-Gln-Met-Gly-Ala-Thr-Ala-Ile-Lys-Gln-Val-Lys-Lys-Leu-Phe-Lys-Lys-Trp-Gly-Trp (seq. id. no. 2).
- 6. The expression system of claim 1, characterized in that said promoter has a DNA sequence that is essentially similar to the promoter elements shown in Figure 4, essentially similar being defined by the presence of direct repeats, TATA boxes and a characteristic spacing between these elements and by the fact that expression initiated at this promoter can be induced by a mechanism similar to the mechanism for the induction of expression of the genes shown in Figure 1.
- 7. The expression system of claim 6, wherein said promoter has the DNA sequence of one of the promoter elements shown in Figure 4.

Replaced by Article 34

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- 8. The expression system of claims 1-7, characterized in that one or more of the said genes are selected from the group of the genes denoted IF, K, R, P, I, T, A in Fig. 1, or the gene/genes are analogous of these genes, analogous genes being defined as genes that bear sequence homology with IF or K or R or P or I or T or A, are isolated from lactic acid bacteria, and are involved in bacteriocin production and/or the regulation thereof in those bacteria, and the said peptide is one of the peptides described in claims 3-5, and that the said promoter is one of the promoters described in claims 6 and 7.
- 9. The expression system of claim 1, characterized in that the said genes are at least one or more genes selected from the group the genes denoted IF,/K, R, T, A, P, and I in Fig. 1, the said promoter has the DNA sequence of one of the promoters according to claim 6 and 7 and the said peptide is Met-Ala-Gly-Asn-Ser-Ser-Asn-Phe-Ile-His-Lys-Ile-Lys-Gln-Ile-Phe-Thr-His-Arg (seq. id. no. 1), or any combination of two or more of the above genes.
 - characterized in comprising any possible combination of genes and promoter elements that are part of claim 1-9, preferably in that it contains a promoter element with the DNA sequence of one of the promoter elements described in claims 6 and 7 operably linked to a gene encoding a desired protein of interest.
- characterized in that it is transformed with the recombinant vector of claim
 10 and contains any possible combination of genes and promoter elements
 that are part of claims 1-9 integrated in the chromosome, and/or in that also
 integrated in its chromosome is a promoter element with the DNA sequence
 of one of the promoter elements described in claims 3 and 4 operably linked
 to an also integrated gene encoding a desired protein of interest.
- 12. The host cells of claim 11, characterized in that some of the said genes and promoter elements are present in plasmids and some are present in the chromosome.
 - 13. The host cells of claims 11-12,

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characterized in that the host is a Gram-positive bacterium, preferably a lactic acid bacterium.

- 14. The host cell of claims 11-13, characterized in that said host possesses the food consumption classification of GRAS (Generally Regarded As Safe).
- characterized in that it is selected from the group consisting of members of the genera Lactobacillus, Lactococcus, Pediococcus, preferably members of the genus Lactobacillus, more preferably of Lactobacillus sake and Lactobacillus plantarum, most preferably of Lactobacillus sake LTH673 and Lactobacillus plantarum C11.
 - 16. Purified peptide, characterized in that it has the amino acid sequence of Met-Ala-Gly-Asn-Ser-Ser-Asn-Phe-Ile-His-Lys-Ile-Lys-Gln-Ile-Phe-Thr-His-Arg (seq. id. no. 1).
- 17. Purified protein,

 characterized in that it is produced by any of the host cells of claims 11-15

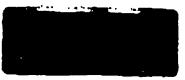
 after or not addition of any of the peptides of claims 3-5.
 - 18. Use the gene expression system according to claim 1-9, in any of the host cells described in claims 1/1-15 to induce gene expression by adding any of the peptides described in claims 3-5.
 - 19. Use of any of the host cells of claims 11-15 in fermentations.
 - 20. Use of any of the host cells of claims 11-15 to produce a desired protein of interest.
- 21. A kit for using the expression system according to claim 1, in lactid acid bacteria, characterized in consisting of:
 - 1) One or more recombinant vectors each vector containing a promotor element identical or similar to one of the promoter elements depicted in Fig. 4, directly followed by a multiple cloning site; these vectors may also contain one or more genes selected from the group K, R, IF, T, A (Fig. 1) or functional analogues of these genes,

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2) Lactic acid bacteria that can function as host strain for these recombinant vectors, and that, depending on the recombinant vector used, may contain one or more genes selected from the group K, R, IF, T, A (Fig. 1) (or functional analogues of these genes) integrated in the chromosome,

3) A peptide that is capable of inducing the expression of genes under control of promoter elements similar or identical to the promoter elements depicted in Fig. 4 and that needs a two component system similar or identical to that encoded by genes K and R (Fig. 1) to exert its inducing action.



CLAIMS

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- Gene expression system. characterized in that it comprises a gene/genes of interest that by genetic engineering have been operably linked to a strongly regulated promoter whose activity can be induced by an unmodified peptide, wherein said 5 promoter and peptide are functional equivalent to promoters and peptides involved in the production of bacteriocins, except nisin, in lactic acid bacteria, and in that the products of two regulatory genes encoding a so called two-component regulatory system are essential for transducing the signal provided by said peptide into a change in activity of said strongly 10 regulated promoter, and in that in naturally occuring lactic acid bacteria said regulatory genes are co-transcribed or closely associated with genes encoding said peptide, wherein the said peptide is a functional analogue of the peptide having the sequences shown in Seq. id. No. 1 and Seq. id. No. 2, and in that said gene/genes of interest are not identical to the genes that are operably 15 linked to said promoter elements in the lactic acid bacterium from which said promoter elements are derived.
- 2. Gene expression system according to claim 1, characterized in that said peptide is capable of inducing its own production and/or the production of one or more bacteriocins in lactic acid bacteria.
 - 3. Gene expression system according to claims 1-2, characterized in that said peptide is identical to the peptide having the sequences of Seq. id. No. 1 and Seq. id. No. 2.
- 4. Gene expression system according to claims 1-3,
 characterized in that said promoter is identical or functionally analogous to
 the promoter elements shown in Fig. 4.
- 5. A recombinant vector,
 characterized in that it comprises the gene/genes operably linked to the
 promoter elements according to claim 1-4, wherein this gene/these genes are
 not identical to the genes that are operably linked to said promoter elements
 in the lactic acid bacterium from which said promoter elements are derived.

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6. A host cell.

characterized in that it contains the gene/genes of claim 5 operably linked to the promoter, and in that the expression of the said gene/genes can be regulated by adding a peptide according to claims 1-4.

- 5 The host cells of claim 6. characterized in that some of the said genes and promoter elements are present in plasmids and some are present in the chromosome.
- The host cells of claims 6-7, characterized in that the host is a Gram-positive bacterium, preferably a 10 lactic acid bacterium.
 - The host cell of claims 6-8, characterized in that said host possesses the food consumption classification of GRAS (Generally Regarded As Safe).
 - 10. The host cells of claims 6-9,
- characterized in that it is selected from the group consisting of members of 15 the genera Lactobacillus, Lactococcus, Pediococcus, profesably members of the genus Lactobacillus, more preferably of Lactobacillus sake and Lactobacillus plantarum, most prefarably of Lactobacillus sake LTH673 and Lactobacillus plantarum C11.
- 20 11. Peptide, characterized in that it has the amino acid sequence of Met-Ala-Gly-Asn-Ser-Ser-Asn-Phe-Ile-His-Lys-Ile-Lys-Gln-Ile-Phe-Thr-His-Arg (seq. id. no. 1).
- 12. Use the gene expression system according to claim 1-4, in any of the host cells described in claims 6-10 to induce gene expression by adding any of the peptides described in claim 3. 25
 - 13. Use of any of the host cells of claims 6-10 in fermentations.
 - 14. Use of any of the host cells of claims 6-10 to produce a desired protein of interest.

AMENDED SHEET

- 15. A kit for using the expression system according to claim 1, in lactid acid bacteria, characterized in consisting of:
- 1) One or more recombinant vectors each vector containing a promotor element identical or similar to one of the promoter elements depicted in Fig. 4, directly followed by a multiple cloning site; these vectors may also contain one or more genes selected from the group K, R, IF, T, A (Fig. 1) or functional analogues of these genes,
- 2) Lactic acid bacteria that can function as host strain for these recombinant vectors, and that, depending on the recombinant vector used, may contain one or more genes selected from the group K, R. IF, T. A (Fig. 1) (or functional analogues of these genes) integrated in the chromosome, such that at least the genes K and R or functional analogues thereof are present in said lactic acid bacteria containing said recombinant vector.

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3) A peptide that is capable of inducing the expression of genes under control of promoter elements similar or identical to the promoter elements depicted in Fig. 4 and that needs a two component system similar or identical to that encoded by genes K and R (Fig. 1) to exert its inducing action.

AMENDED SHEET